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(54) **METHOD FOR STIMULATING THE IMMUNE SYSTEM**

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WO	WO 95/00103	1/1995
WO	WO 96/02143	2/1996
WO	WO 96/23065	8/1996
WO	WO 97/39120	10/1997
WO	WO 99/50411	10/1999

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(22) Filed: **Sep. 7, 2007**

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Jul. 25, 1998	(EP)	.....	98113974

(51) **Int. Cl.**

<b>A01N 43/04</b>	(2006.01)
<b>C12Q 1/68</b>	(2006.01)
<b>C12P 19/34</b>	(2006.01)
<b>C07H 21/02</b>	(2006.01)
<b>C07H 21/04</b>	(2006.01)

(52) **U.S. Cl.**

USPC ..... **514/44**; 435/6; 435/91.1; 435/91.31;  
435/455; 514/1; 514/2; 536/23.1; 536/24.31;  
536/24.5

(58) **Field of Classification Search**

USPC ..... 435/6, 91.1, 91.31, 375, 455; 514/1, 2,  
514/44; 536/23.1, 24.5, 23.5, 24.31  
See application file for complete search history.

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(57) **ABSTRACT**

Medicament comprising a combination of

at least one inhibitor of the effect of a substance negatively effecting an immune response, the substance selected from the group consisting of TGF- $\beta$  and its receptors, VEGF and its receptors, interleukin 10 (IL-10) and its receptors, PGE<sub>2</sub> and its receptors, wherein the inhibitor has a molecular weight of less than 100 kDa and

at least one stimulator positively effecting an immune response.

**6 Claims, 11 Drawing Sheets**

1.	TGF-B2-1	C ACA CAG TAG TGC A
2.	TGF-B2-2	GC ACA CAG TAG TGC
3.	TGF-B2-3	GC TTG CTC AGG ATC TGC
4.	TGF-B2-4	TAC TCT TCG TCG CT
5.	TGF-B2-5	C TTG GCG TAG TAC T
6.	TGF-B2-6	G TAA ACC TCC TTG G
7.	TGF-B2-7	GT CTA TTT TGT AAA CCT CC
8.	TGF-B2-8	GC ATG TCT ATT TTG TAA ACC
9.	TGF-B2-9	CGG CAT GTC TAT TTT GTA
10.	TGF-B2-10	G GCA TCA AGG TAC C
11.	TGF-B2-11	CTG TAG AAA GTG GG
12.	TGF-B2-12	AC AAT TCT GAA GTA GGG T
13.	TGF-B2-13	T CAC CAA ATT GGA AGC AT
14.	TGF-B2-14	GCT TTC ACC AAA TTG GAA GC
15.	TGF-B2-15	CTG GCT TTT GGG TT
16.	TGF-B2-16	T CTG ATA TAG CTC AAT CC
17.	TGF-B2-17	T CCT AGT GGA CTT TAT AG
18.	TGF-B2-18	T TTT TCC TAG TGG ACT
19.	TGF-B2-19	C AAT TAT CCT GCA CAT TTC
20.	TGF-B2-20	GC AAT TAT CCT GCA CA
21.	TGF-B2-21	GC AGC AAT TAT CCT GC
22.	TGF-B2-22	TG GCA TTG TAC CCT
23.	TGF-B2-23	TG TGC TGA GTG TCT
24.	TGF-B2-24	CC TGC TGT GCT GAG TG
25.	TGF-B2-25	C TTG GGT GTT TTG C
26.	TGF-B2-26	T TTA GCT GCA TTT GCA AG
27.	TGF-B2-27	G CCA CTT TTC CAA G
28.	TGF-B2-14/1	CTT TCA CCA AAT TGG AAG
29.	TGF-B2-14/2	CAC CAA ATT GGA AGC
30.	TGF-B2-14/3	TCA CCA AAT TGG AAG C
31.	TGF-B2-15/1	CTC TGG CTT TTG GG
32.	TGF-B2-9/1	CGG CAT GTC TAT TTT G
33.	TGF-B1-1	CGA TAG TCT TGC AG
34.	TGF-B1-2	GTC GAT AGT CTT GC
35.	TGF-B1-3	CTT GGA CAG GAT CT
36.	TGF-B1-4	CCA GGA ATT GTT GC
37.	TGF-B1-5	CCT CAA TTT CCC CT
38.	TGF-B1-6	GAT GTC CAC TTG CA
39.	TGF-B1-7	CTC CAA ATG TAG GG
40.	TGF-B1-8	ACC TTG CTG TAC TG
41.	TGF-B1-9	GTA GTA CAC GAT GG
42.	TGF-B1-10	CAC GTA GTA CAC GA
43.	TGF-B1-11	CAT GTT GGA CAG CT
44.	TGF-B1-12	GCA CGA TCA TGT TG
45.	TGF-B1-13	TGT ACT CTG CTT GAA C
46.	TGF-B1-14	CTG ATG TGT TGA AGA ACA
47.	TGF-B1-15	CTC TGA TGT GTT GAA G
48.	TGF-B1-16	GGA AGT CAA TGT ACA G
49.	TGF-B1-17	CAT GTC GAT AGT CTT GCA
50.	TGF-B1-18	AGC TGA AGC AAT AGT TGG
51.	TGF-B1-19	GTC ATA GAT TTC GTT GTG
52.	TGF-B1-20	CTC CAC TTT TAA CTT GAG
53.	TGF-B1-21	TGC TGT ATT TCT GGT ACA
54.	TGF-B1-137	CGA TAG TCT TGC AG

Fig. 1-1

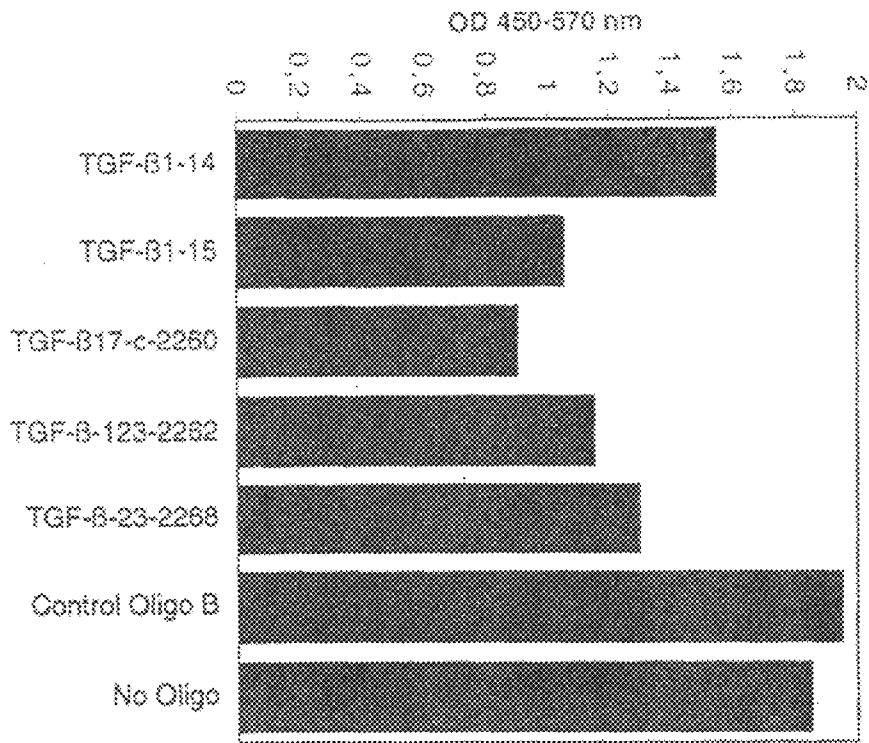
55.	b1-N17	TCC TCT TCG ACT GCT CTC
56.	b2-N14	CGA AGG TTA AAC CAC TTT CG
57.	b2-N24	GTG AGT CGT GTC GTC C
58.	TGF-β2-98-1	CATCGTTGTCGTCG
59.	TGF-β2-98-2	CGCTTCTTCCGCCG
60.	TGF-β2-98-3	CGAAGGAGAGCCATTCG
61.	TGF-β2-98-4	CGATGTAGCG
62.	TGF-β2-98-5	CGTCAAATCG
63.	TGF-β2-98-6	CGTAGTACTCTTCGTCG
64.	TGF-β2-98-7	CGCGCTCGCAGGCG
65.	TGF-β2-98-8	CGGCCGCCCTCCGGCTCG
66.	TGF-β2-98-9	CGCGGATCGCCTCG
67.	TGF-β2-98-10	GAGCGCGACCGTGAC
68.	TGF-β-17-c-2260	ACC TCC TTG GCG TAG TA
69.	TGF-β-12-9/20-2261	AGG GCG GCA TGT CTA TTT TG
70.	TGF-β-123-2262	CAG AAG TTG GCA TTG TAC
71.	TGF-β-12-9/22-2263	AGG GCG GCA TGT CTA TTT TGT A
72.	TGF-β-23-2268	TGG GAC ACG CAG CAA GG
73.	TGF-β1-98-1	CGGGGGCGGGCGGGG
74.	TGF-β1-98-2	CGGGGGCGGGCGGGGCG
75.	TGF-β1-98-3	CGGGCGCCGCCGAGGCGCCCG
76.	TGF-β1-98-4	CCGAGGTCCTTGCGG
77.	TGF-β1-98-5	CGGCGGTGCCGGGA
78.	TGF-β1-98-6	CTCGGCGGCCGCTAG
79.	TGF-β1-98-7	CGCTAAGGCG
80.	TGF-β1-98-8	CCGCACAACCTCCGG
81.	TGF-β1-98-9	GCGAGTCGCTGG
82.	TGF-β1-98-10	CGGTTGCTGAGGTATCG
83.	TGF-β1-98-11	CCGGGAGAGCAACACGG
84.	TGF-β1-98-12	CGCTTCTCG
85.	TGF-β1-98-13	CCATTAGCACGCGGG
86.	TGF-β1-98-14	CGGGCTCCG
87.	TGF-β1-98-15	CCGGCCACCCGGTCGCGG
88.	TGF-β1-98-16	CGAGCACGGCCTCG
89.	TGF-β1-98-17	CGGGCAGCGGGCCGGGCG
90.	TGF-β1-98-18	CGCGGATGGCCTCG
91.	TGF-β1-98-19	CGATGCGCTTCCG
92.	TGF-β1-98-20	CCC CGCGCCGGCGGG
93.	TGF-β1-98-21	CGCAGCCCCGAGGGCG
94.	TGF-β1-98-22	CGGCGCCCCCGG
95.	TGF-β1-98-23	CGGCACTGCCGAGAGCGCG
96.	TGF-β1-98-24	CGGGGATGAAGGCGGGC
97.	TGF-β1-98-25	CGGGTCGGCGACTCCCCG
98.	TGF-β1-98-26	CGCCTGAGGGACGCCG
99.	TGF-β1-98-27	AAGCGTCCCCGGCG
100.	TGF-β1-98-28	CGCGGGGACGCGTCGCG
101.	TGF-β1-98-29	CCCCGGCCCTCCGG
102.	TGF-β1-98-30	CGGCGGCGGCTCG
103.	TGF-β1-98-31	CGCTCCGGGCGGAGGCCG
104.	TGF-β1-98-32	CGGCCCCGCGGGCG
105.	TGF-β1-98-33	CGGACGGGGCGTCC
106.	TGF-β1-98-34	CGGCCGGGGCCCTCG

Fig. 1-2

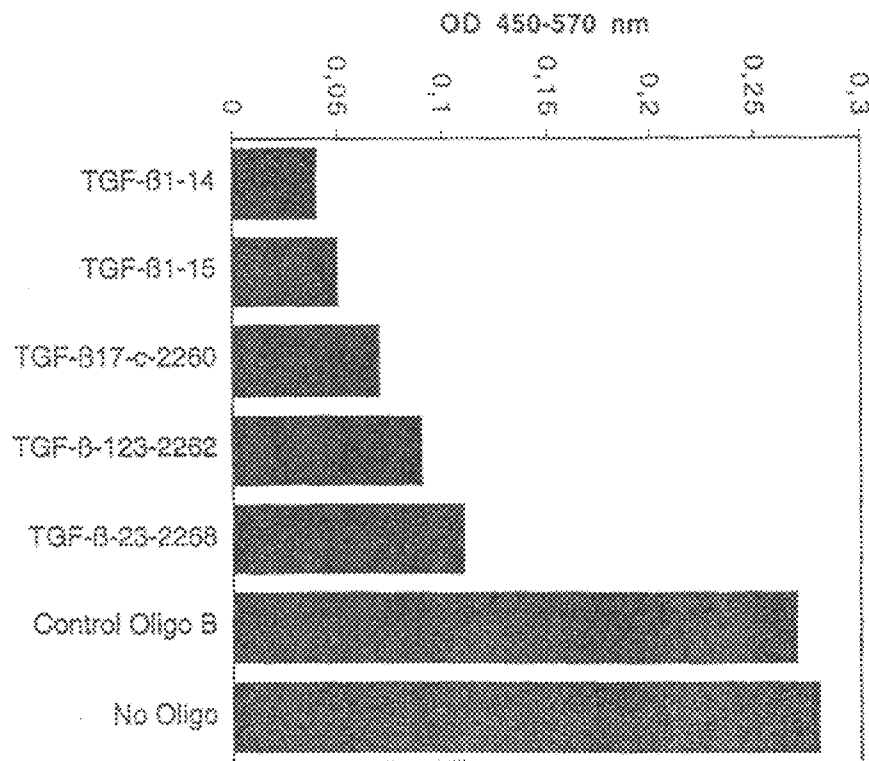
107.	TGF-β3-98-1	TCGAGCTTCCCCGA
108.	TGF-β3-98-2	CCCGGAGCCGAAGG
109.	TGF-β3-98-3	CCCGAGGAGCGGG
110.	TGF-β3-98-4	ACGCAGCAAGGCGA
111.	TGF-β3-98-5	CGGGTTGTTCGAGCCG
112.	TGF-β3-98-6	CGGCAGTGCCCCG
113.	TGF-β3-98-7	CGGAATTCTGCTCG
114.	TGF-β3-98-8	TTCGTTGTGCTCCG
115.	TGF-β3-98-9	ATCCGACTCGGTG
116.	TGF-β3-98-10	ACGTGGGTCATCACCGT
117.	TGF-β3-98-11	CGAAGAAGCG
118.	TGF-β3-312	CCT AAT GGC TTC CA
119.	VEGF-98-1	CGGCCGCGGTGTGT
120.	VEGF-98-2	CGGGAATGCTTCCGCCG
121.	VEGF-98-3	CGGCTCACCGCCTCGGC
122.	VEGF-98-4	CACGTCTGCGGATC
123.	VEGF-98-5	CCCCGCATCGCATCAGGG
124.	VEGF-98-6	CGCCTTGCAACGCG
125.	VEGF-98-7	CCGACCGGGGCCGG
126.	VEGF-49	GTTTCATGGTTTCGG
127.	VEGF-55	GCAGAAAGTTCATGG
128.	VEGF-188	GCTGATAGACATCC
129.	VEGF-190	GCGCTGATAGACAT
130.	VEGF-194	GTAGCTGCGCTGATAG
131.	VEGF-253	CTCGATCTCATCAG
132.	VEGF-255	ATGTA CTGATCTCATC
133.	VEGF-260	GAAGATGTA CTGATC
134.	VEGF-263	CTTGAAGATGTA CTG
135.	VEGF-292	GCATCGCATCAGGG
136.	VEGF-294	CCGCATCGCATCAG
137.	VEGF-422	CATTTGTTGTGCTGTAGG
138.	VEGF-434	GGTCTGCATTCACATTTG
139.	VEGF-441	CTTTGGTCTGCATTC
140.	VEGF-445	CTTTCTTTGGTCTGC
141.	VEGF-450	GCTCTATCTTTCTTTGG
142.	VEGF-455	GTCTTGCTCTATCTTTC
143.	VEGF-459	CTTGTCTTGCTCTATC
144.	VEGF-596	CATCTGCAAGTACGTTTCG
145.	VEGF-598	CACATCTGCAAGTACGTT
146.	VEGF-600	GTCACATCTGCAAGTACG
147.	VEGF-600-2	CATCTGCAAGTACG
148.	VEGF-601	CACATCTGCAAGTAC
149.	VEGF-604	GTCACATCTGCAAG
150.	VEGF-607	CTTGTACATCTGC
151.	VEGF-607-2	GGCTTGTCACATCTGC
152.	VEGF-610	CTCGGCTTGTCACATC
153.	VEGF-638	CTCCTTCCTCCTGC
154.	VEGF-766	GCT TGA AGA TGT ACCT CG
155.	VEGF-r-1062	CGT TGC TCT CCG ACG
156.	flt-1165	GAC ACG GCC TTT TCG
157.	flt-rm-2115	CCA GCA GCT GAC CAT GG
158.	flkl1/kdr-m-2315	GAA ATC GAC CCT CGG
159.	MCP-1-Rec-A/B-571	GCA TGT TGT GGA TG
160.	MCP-1-1954	GCA GAG ACT TTC ATG C
161.	MCP-1-1955	ATA ACA GCA GGT GAC TGG

Fig. 1-3

162.	MCP-1-1956	GAA CCC ACT TCT GC
163.	MCP-1-2761	GAC ACT TGC TGC TG
164.	MCP-1-2762	CCA CTT CTG CTT GGG
165.	VEGF-703	CTG CAA GTA CGT TCG
166.	flt-1567	TCC CTT ATG ATG CCA GCA AGT G
167.	TGF- $\beta$ -Rec-I-796	CCA GCA ATG ACA GC
168.	TGF- $\beta$ -1-rwk-1	G GGA AAG CTG AGG C
169.	TGF- $\beta$ -1-rwk-2	T CGA GGG AAA GCT GA
170.	TGF- $\beta$ -1-rwk-3	C CTC GAG GGA AAG C
171.	TGF- $\beta$ -1-rwk-4	GG GCT GGT GTG GTG
172.	TGF- $\beta$ -1-rwk-5	GA ACA GGG CTG GTG TG
173.	TGF- $\beta$ -1-rwk-6	G AAC AGG GCT GGT G
174.	TGF- $\beta$ -1-rwk-7	AG AGC GCG AAC AGG
175.	TGF- $\beta$ -1-rwk-8	GA GAG CGC GAA CAG G
176.	TGF- $\beta$ -1-rwk-9	CGA GAG CGC GAA CAG
177.	TGF- $\beta$ -1-rwk-10	CCC CTG GCT CGG GGG
178.	TGF- $\beta$ -1-rwk-11	C CCT GGC TCG GGG
179.	TGF- $\beta$ -1-rwk-12	C CCC TGG CTC GGG G
180.	TGF- $\beta$ -1-rwk-13	TCC CCC TGG CTC GG
181.	TGF- $\beta$ -1-rwk-14	C TCC CCC TGG CTC G
182.	TGF- $\beta$ -1-rwk-15	TGC GCT TCC GCT TCA C
183.	TGF- $\beta$ -1-rwk-16	CC TCG ATG CGC TTC
184.	TGF- $\beta$ -1-rwk-17	G ATG GCC TCG ATG C
185.	TGF- $\beta$ -1-rwk-18	G GAT GGC CTC GAT GC
186.	TGF- $\beta$ -1-rwk-19	ATG GCC TCG ATG CGC TT
187.	TGF- $\beta$ -3-rwk-1	TC AGC AGG GCC AGG
188.	TGF- $\beta$ -3-rwk-2	GCA AAG TTC AGC AGG GC
189.	TGF- $\beta$ -3-rwk-3	GG CAA AGT TCA GCA GG
190.	TGF- $\beta$ -3-rwk-4	GT GGC AAA GTT CAG CAG G
191.	TGF- $\beta$ -3-rwk-5	GTG GCA AAG TTC AG
192.	TGF- $\beta$ -3-rwk-6	GAC CGT GGC AAA GTT CAG
193.	TGF- $\beta$ -3-rwk-7	AGA GAG GCT GAC CGT
194.	TGF- $\beta$ -3-rwk-8	GAC AGA GAG AGG CTG AC
195.	TGF- $\beta$ -3-rwk-9	A CAG AGA GAG GCT GA
196.	TGF- $\beta$ -3-rwk-10	GT GGA CAG AGA GAG G
197.	TGF- $\beta$ -3-rwk-11	CA AGT GGA CAG AGA GAG G
198.	TGF- $\beta$ -3-rwk-12	TCT TCT TGA TGT GGC C
199.	TGF- $\beta$ -3-rwk-13	CC CTC TTC TTC TTG ATG
200.	TGF- $\beta$ -3-rwk-14	C ACC CTC TTC TTC T
201.	TGF- $\beta$ -3-rwk-15	A TGG ATT TCT TTG GCA T
202.	TGF- $\beta$ -3-rwk-16	GGA TTT CTT TGG C
203.	TGF- $\beta$ -3-rwk-17	AA GTT GGA CTC TCT TCT C
204.	TGF- $\beta$ -3-rwk-18	TAA GTT GGA CTC TCT TCT
205.	TGF- $\beta$ -3-rwk-19	GAC CTA AGT TGG ACT C
206.	TGF- $\beta$ -3-rwk-20	T TTC TAG ACC TAA GTT GG
207.	TGF- $\beta$ -3-rwk-21	CT GAT TTC TAG ACC TAA G
208.	TGF- $\beta$ -3-rwk-22	G AAG CAG TAA TTG GTG T
209.	TGF- $\beta$ -3-rwk-23	GG AAT CAT CAT GAG G
210.	TGF- $\beta$ -3-rwk-24	GGG AAT CAT CAT GAG
211.	TGF- $\beta$ -3-rwk-25	G GTT GTC GAG CCG GT
212.	TGF- $\beta$ -3-rwk-26	GTC CTC CCA ACA TAG TA
213.	TGF- $\beta$ -3-rwk-27	GG GTC CTC CCA ACA

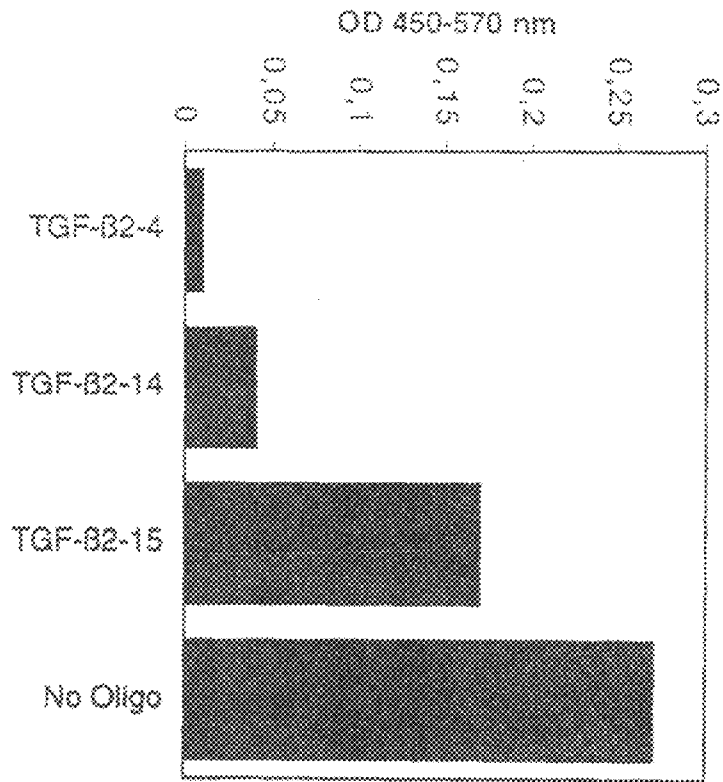


A.

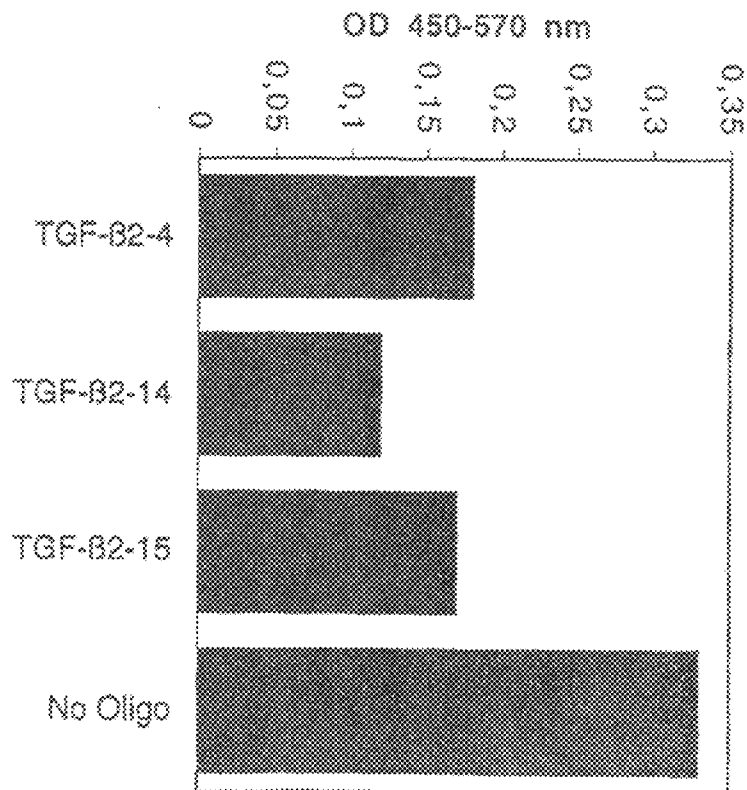


B.

Figure 2



A.



B.

Figure 3

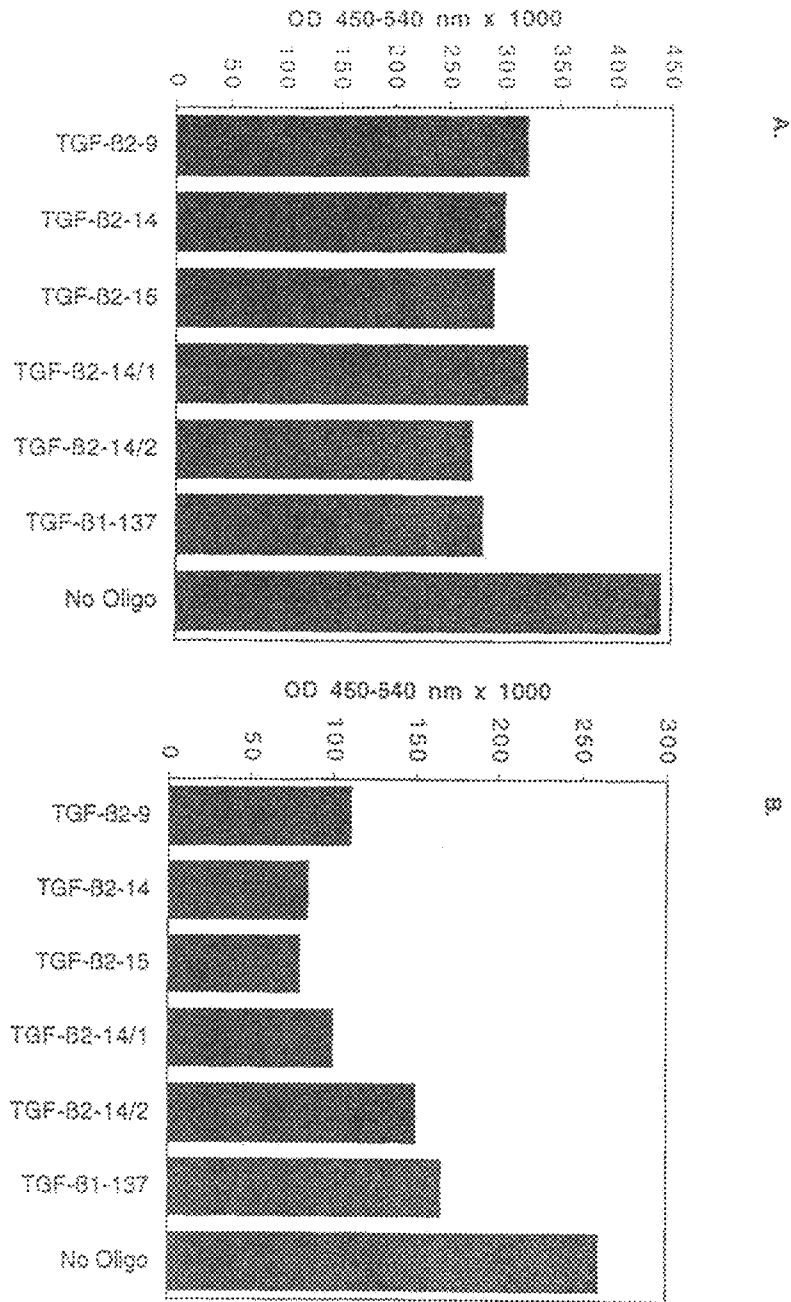


Figure 4

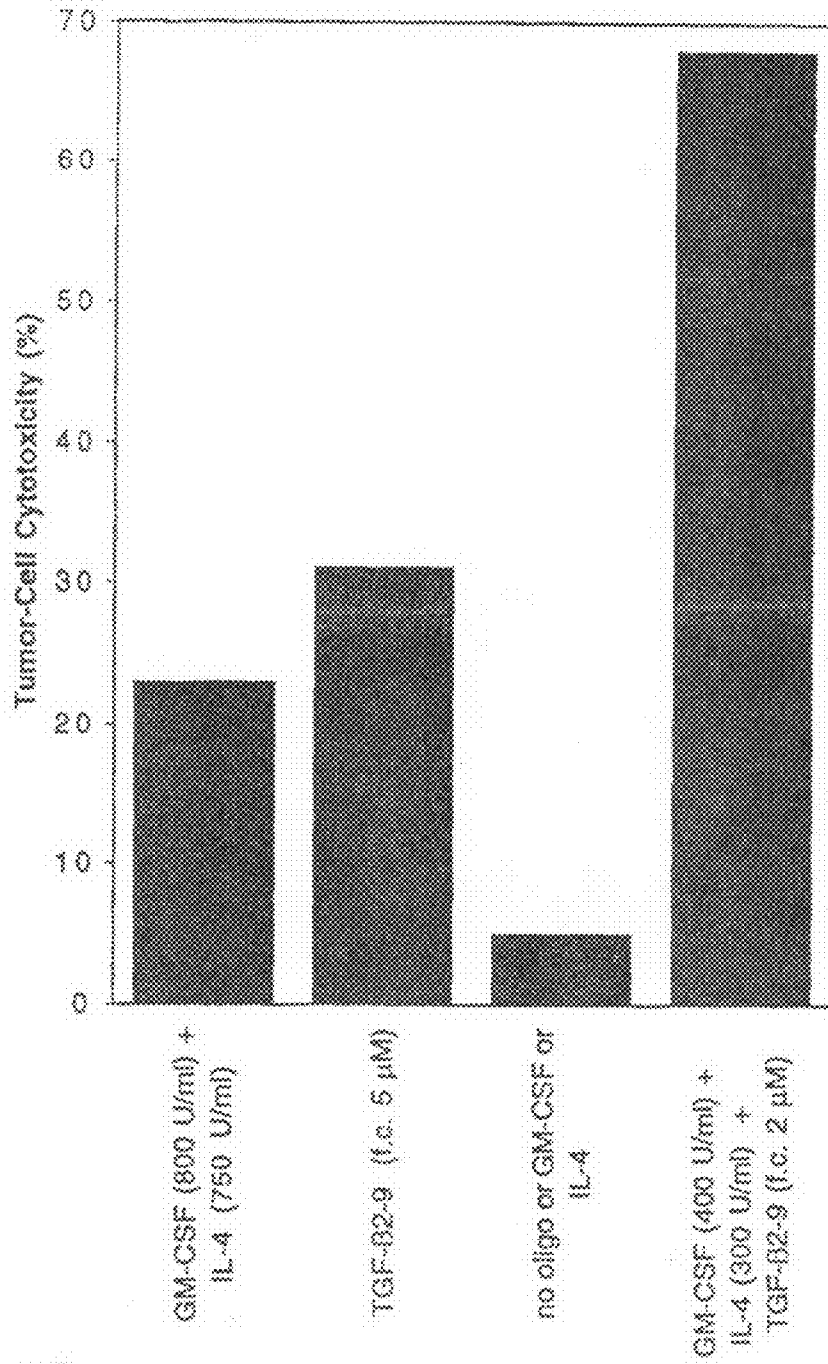
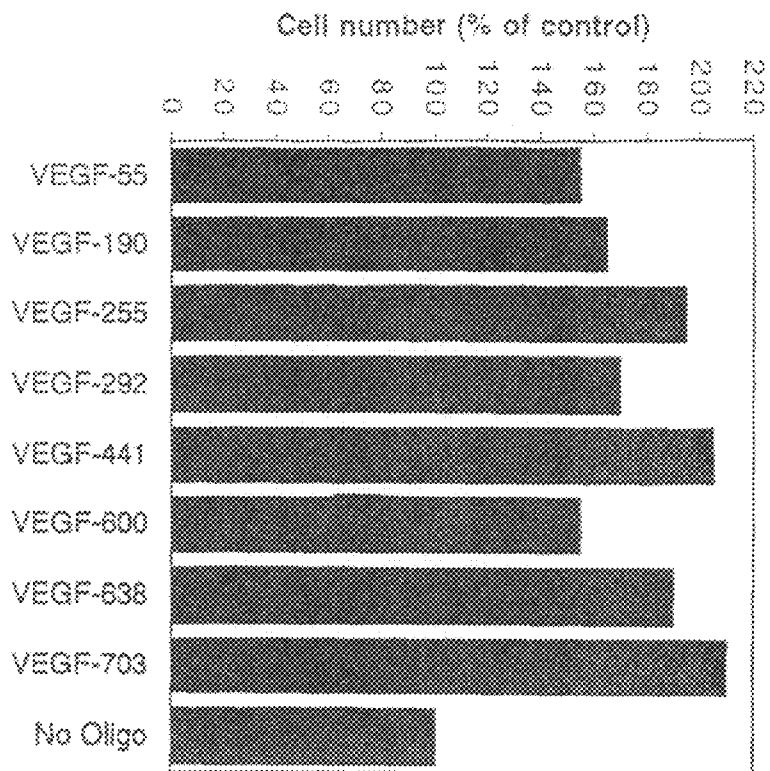
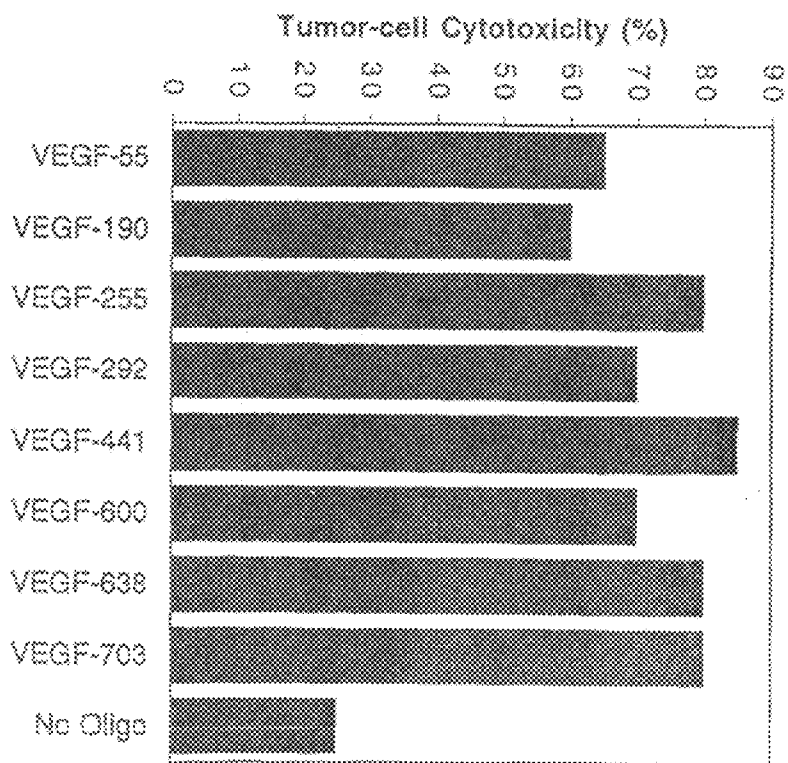


Figure 5

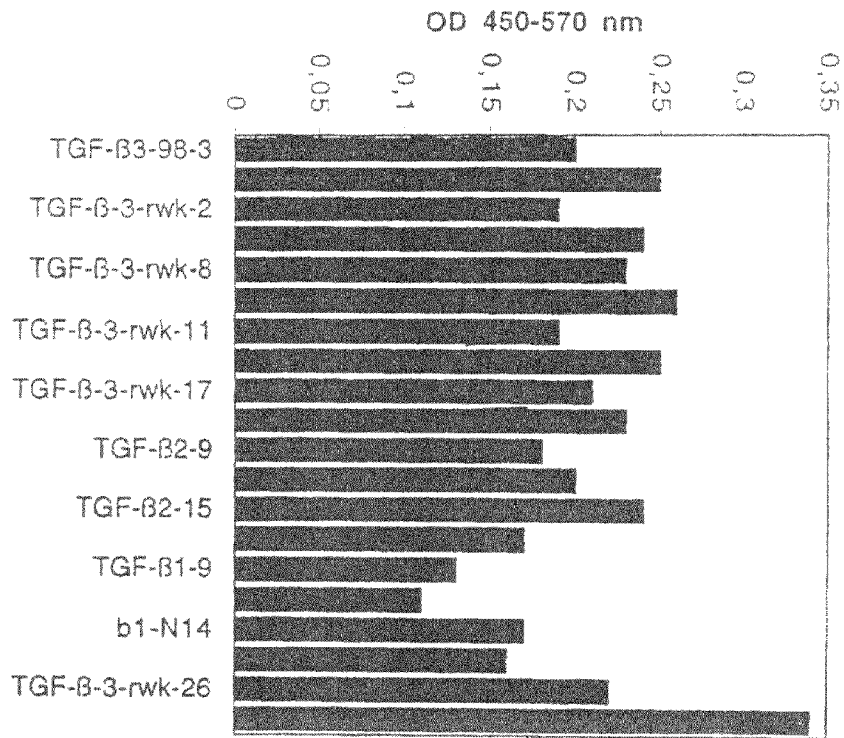


A.

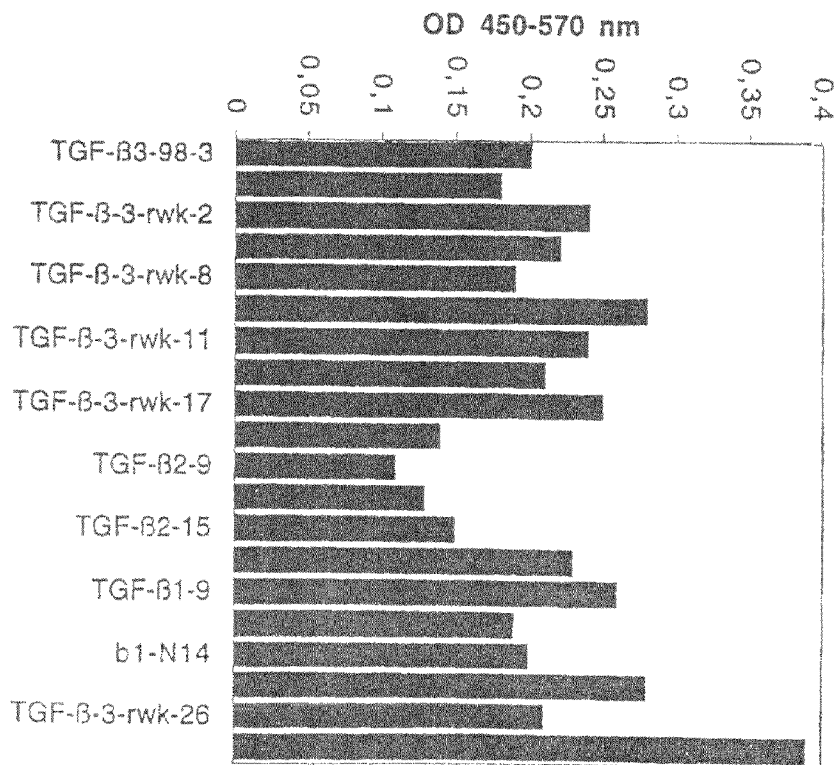


B.

Figure 6



A.



B.

Figure 7

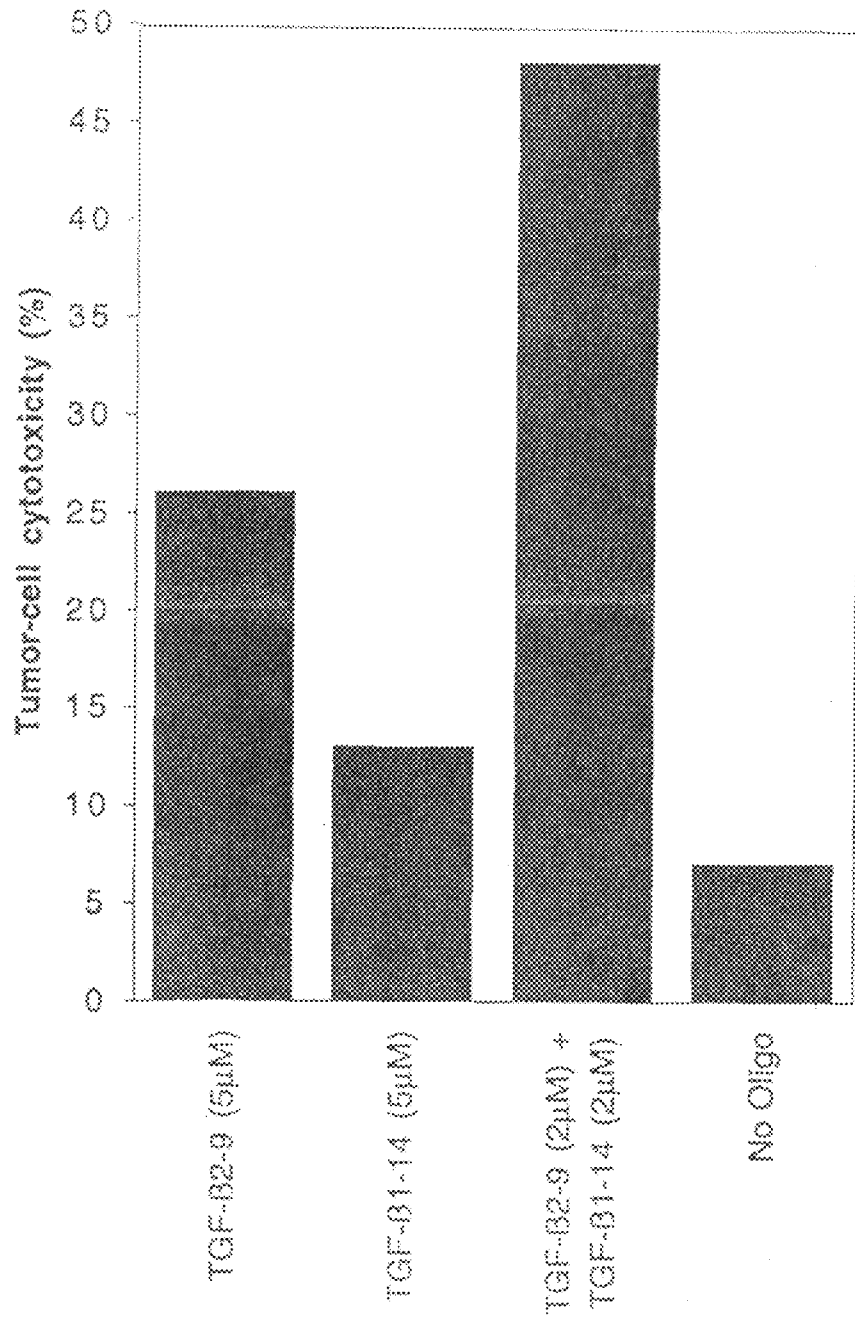


Figure 8

**METHOD FOR STIMULATING THE IMMUNE SYSTEM**

Two different approaches have been used in the prior art to enhance the immune response against neoplastic cells. One approach uses the addition of cytokines like interleukin-2 (IL-2) or transfection of tumor cells and/or immune cells with genes coding for cytokines like IL-2 or other proteins enhancing the immune response like transfection of tumor cells with lymphotactin or like transfection of T-lymphocytes with CD-40 Ligand.

The second approach uses the inhibition of immunosuppressive molecules to enhance the body's immune response to tumor cells. Thus, J. NEUROSURG. 78 (1993) 944-51, Jachimczak et al. (1993) and WO 94/25588, Schlingensiepen et al. (1994) teach the use of antisense oligonucleotides targeted to TGF- $\beta$  to reverse tumor-induced immunosuppression.

Several documents in the prior art teach that a combination of these two approaches is either not efficacious or is not beneficial over use of one of the two approaches used alone.

Thus, CANCER BIOTHER. 8(2), 1993, 159-170, Gridley et al., as well as CANCER BIOTHER. 9(4), 1994, 317-327, Mao et al., both teach that a combination of anti-transforming growth factor-beta antibody with IL-2 does not cause significant antitumor effects.

Furthermore, PROC. NATL. ACAD. SCI 93, (1996), 2909-2914, Fakhrai et al., teaches that a combination of transfection with genes encoding antisense sequences to transforming growth factor beta (TGF- $\beta$ ) TGF- $\beta$  mRNA with transfection of IL-2 into tumor cells does not increase the immune response against the tumor compared to transfection with TGF- $\beta$  antisense alone.

Surprisingly, in contrast, certain combinations of stimulators and inhibitors are more efficacious than either approach alone.

The present invention discloses a medicament comprising a combination of

at least one inhibitor of the effect of a substance negatively effecting an immune response, the substance selected from the group consisting of TGF- $\beta$  and its receptors, VEGF and its receptors, interleukin 10 (IL-10) and its receptors, PGE<sub>2</sub> and its receptors, wherein the inhibitor has a molecular weight of less than 100 kDa and at least one stimulator positively effecting an immune response.

In a preferred embodiment, the inhibitor is inhibiting the synthesis or function of molecules suppressing or downregulating or negatively affecting the immune response. The inhibitor can be an oligonucleotide which may function as an antisense nucleotide or a ribozyme or it may be an antibody fragment derived from an anti-body e.g. a fab-fragment or a single chain antibody.

Preferably, the stimulator is positively effecting the immune response by increasing presentation of antigens and/or enhancing proliferation and/or function of immune cells.

In a preferred embodiment, the stimulator is enhancing the synthesis or function of molecules stimulating, enhancing, upregulating and/or positively regulating the immune response. In particular, the stimulator is stimulating and/or enhancing the synthesis and/or the function of factors such as GM-CSF, SCF, CSF, IFN- $\gamma$ , FLT-3-ligand as well as monocyte chemotactic proteins (MCP-1), interleukin-2, interleukin-4, interleukin-12 and/or interleukin-18 or is one of the mentioned interleukins or is selected from the group consisting of viruses, viral antigens, antigens expressed in tumor cells or pathogens but not in normal cells, organspecific antigens

expressed in affected organs which are not essential for the organism, e.g. prostate, ovary, breast, melanine producing cells.

The stimulators are preferably selected from

- a) Chemokines, including lymphotactin and/or immune cell attracting substances and/or
- b) viruses and/or parts of viruses, including retroviruses, adenoviruses, papillomaviruses, Epstein-Barr-Viruses, Viruses that are non-pathogenic including Newcastle-Disease virus, Cow-pox-virus and/or
- c) autologous and/or heterologous MHC-Molecules and/or
- d) molecules involved in antigen processing and/or
- e) molecules involved in antigen presentation and/or
- f) molecules involved in mediating immune cell effects and/or
- g) molecules involved in mediating immune cell cytotoxic effects and/or
- h) molecules involved in antigen transportation and/or
- i) co-stimulatory molecules
- j) peptides enhancing recognition by immune cells and/or cytotoxic effects of immune cells
- k) the peptides containing one or more amino acids differing between a protein in the target cell from the other cells within an organism
- l) the peptides according to j) being
  - Peptides containing one or more mutations and/or amino acid substitutions of the ras protein amino and/or
  - Peptides containing one or more mutations and/or amino acid substitutions of the p53 protein and/or
  - Peptides containing one or more mutations and/or amino acid substitutions of the EGF-Receptor protein and/or
  - Peptides containing one or more mutations and/or amino acid substitutions of fusion peptides and/or fusion proteins and/or
  - Peptides containing one or more mutations and/or amino acid substitutions and/or amino acid substitutions caused by gene rearrangements and/or gene translocations and/or
  - Peptides containing one or more mutations and/or amino acid substitutions of the retinoblastoma protein and/or
  - Peptides containing one or more mutations and/or amino acid substitutions of proteins coded by oncogenes and/or protooncogenes and/or
  - Peptides containing one or more mutations and/or amino acid substitutions of proteins coded by anti-oncogenes and/or tumor suppressor genes and/or
  - Peptides derived from proteins differing in the target cell by one or more amino acids from the proteins expressed by other cells in the same organism and/or
  - Peptides derived from viral antigens and/or coded by viral nucleic acids and/or
  - Peptides derived from proteins expressed in a diseased organ but not in the nervous system, muscle, hematopoietic system or other organs essential for survival. Diseased organs are e.g. prostate, ovary, breast, melanine producing cells and the like.
- m) tumor cell extracts and/or tumor cell lysates and/or adjuvants,
- n) fusion cells of dendritic and tumor cells.

These fusion cells are hybridoma cells derived from a mixture of dendritic cells and tumor cells. Dendritic cells are generated e.g. by treatment of PBMC with GM-CSF and IL-4 or a mixture of GM-CSF, IL-4 and IFN- $\gamma$  or FLT-3 ligand. Fusion of dendritic cells with tumor cells can be achieved e.g. using PEG (polyethylene glycole) or electrofusion.

Surprisingly, treatment of PBMC with VEGF-oligonucleotides enhanced the number and/or effectiveness of dendritic cells.

In one embodiment of the invention the inhibitor is an oligonucleotide. Preferably the oligonucleotides of FIG. 1 are useful in the medicament of the present invention.

In a further embodiment, the invention provides oligonucleotides having one of the sequences given in FIG. 1-2 to 1-4.

Also oligonucleotides having 1 to 10 additional-nucleotides at the 5'- or 3'-end are part of the invention.

Oligonucleotide sequences used for transfection are usually much longer sequences than those used for antisense oligonucleotides, which usually do not exceed 30 bases in length and are applied as short single-stranded sequences and are not integrated into a vector system.

Since transfected sequences are usually much longer than oligonucleotides, if cross inhibition of different members of a protein family would occur with the antisense technology, such cross inhibition of other mRNAs than the target mRNA, is much more likely with transfected antisense sequences, compared to oligonucleotides. However, Cell Growth Differ, Vol. 6(12), February 1995, pages 1635-1642, Huang, F. et al. teaches "only the K6 transfectant exhibited 39 and 33% respectively of the levels of TGF beta1 mRNA and active secreted TGF beta1 protein of the parental line. K6 exhibited no change in TGF beta2 expression and TGF beta3 expression was not detected in either parental or transfectant cell line."

It was therefore surprising to find oligonucleotides according to this invention, which were able to significantly reduce expression of both, TGF- $\beta_1$  as well as TGF- $\beta_2$  e.g. TGF- $\beta_1$ -14, TGF- $\beta_1$ -15, TGF- $\beta$ -17-c-2260, TGF- $\beta$ -123-2262, TGF- $\beta$ -23-2268, TGF- $\beta$ -2-4, TGF- $\beta$ -2-14, TGF- $\beta$ -2-15, TGF- $\beta$ -2-9, TGF- $\beta$ -2-14/1, TGF- $\beta$ -2-14/2, TGF- $\beta$ -1-136. Furthermore surprisingly oligonucleotides were designed, which were able to significantly reduce expression of TGF- $\beta_2$  as well as TGF- $\beta_3$ .

Surprisingly even oligonucleotides were found, which were able to significantly reduce expression of TGF- $\beta_2$  as well as TGF- $\beta_1$ , and TGF- $\beta_3$ , e.g. b1-N17, b1-N14, b1-N24, TGF- $\beta$ -2-9, TGF- $\beta$ -2-14, TGF- $\beta$ -2-15, TGF- $\beta$ -17-c-2260, TGF- $\beta$ -12-9/20-2261, TGF- $\beta$ -123-2262, TGF- $\beta$ -12-9/22-2263, TGF- $\beta$ -23-2268, TGF- $\beta$ -1-98-11, TGF- $\beta$ -1-98-23, TGF- $\beta$ -3-98-7, TGF- $\beta$ -3-98-10, TGF- $\beta$ -1-rwk-5, TGF- $\beta$ -3-rwk-2, TGF- $\beta$ -1-rwk-5, TGF- $\beta$ -3-rwk-9, TGF- $\beta$ -3-rwk-23, TGF- $\beta$ -1-3, TGF- $\beta$ -1-10.

Thus oligonucleotides which are effective against expression of at least two of TGF- $\beta_1$ , TGF- $\beta_2$  and/or TGF- $\beta_3$  are also part of the invention.

These findings were also surprising in view of the fact that sequence comparison between the mRNAs of TGF- $\beta_2$ , TGF- $\beta_1$ , and TGF- $\beta_3$  showed that not a single sequence of 20 bases in length could be found that would be identical within the three different mRNAs. Even if such a hypothetical sequence had really existed, inhibition of the three mRNAs by such a hypothetical consensus sequence would have been extremely unlikely, since it is well known in the art that only a small minority of antisense sequences complementary to a certain mRNA actually exert a so-called antisense effect, i.e. inhibit expression of the respective protein.

Endothelial synthesis of monocyte chemotactic protein-1 (MCP-1) has been implicated in the regulation of monocyte recruitment for extravascular pools both under physiological and inflammatory conditions.

MCP-1 antisense oligonucleotides were able to modulate monocyte infiltration and were thus anti-inflammatory.

These antisense-oligonucleotides are useful for the treatment of inflammatory diseases e.g. asthma, morbus crohn,

collitis ulcerosa, diabetes, glomerulonephritis, acute respiratory distress syndrome and arteriosclerotic plaque formation.

In a preferred embodiment of the invention the oligonucleotides and/or ribozymes and/or nucleic acids have modifications at the bases, the sugars and/or the phosphate moieties of the oligonucleotides.

In a further preferred embodiment of the invention the oligonucleotides and/or ribozymes and/or nucleic acids have modifications wherein the modifications are phosphorothioate (S-ODN) internucleotide linkages and/or methylphosphonate internucleotide linkages and/or phosphoramidate linkages and/or peptide linkages and/or 2'-O-derivatives, such as 2'-O-methyl or 2'-O-methoxyethoxy modifications of the sugar and/or modifications of the bases.

In a further preferred embodiment of the invention the oligonucleotides and/or ribozymes and/or nucleic acids are coupled to or mixed with folic acid, hormones, steroid hormones such as oestrogene, progesterone, corticosteroids, mineral corticoids, peptides, proteoglycans, glycolipids, phospholipids, polyethylene imine or other poly cations and derivatives therefrom.

Furthermore, the present invention provides a method of treating hyperproliferative diseases, neoplasms or infectious diseases by administering a medicament of the invention to patients in need thereof. The method is especially useful for the treatment of leukemia, non-hodgkin lymphoma, hodgkin lymphoma, bronchial carcinoma, esophageal carcinoma, colorectal carcinoma, gastric carcinomas, intestinal tumors, hepatic tumors, gall bladder and gallduct carcinomas, pancreatic carcinoma, anal carcinoma, breast cancer, ovarian carcinoma, cervical carcinoma, endometrium carcinoma, prostatic carcinoma, bladder carcinoma, malignant melanoma, brain tumors, and sarcomas.

The necessary doses of the medicament of the present invention depend on the disease and the severity of the disease. Whereas higher levels are more effective, they often have a higher degree of side effects. Suitable doses are selected to obtain concentrations of the oligonucleotides in the range of 0.1 to 10  $\mu\text{mol/l}$  and concentrations of the cytokines in the range of 10 to 1.000 U/ml in the patient blood.

In a preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ and the stimulator positively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally).

In another preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally) to the tumor and the stimulator positively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ. In another preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally) to the tumor and the stimulator positively effecting an immune response is applied systemically (e.g. i.v. or s.c. or orally).

In another preferred embodiment of the invention the inhibitor of the effect of a substance negatively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ and the stimulator positively effecting an immune response is applied locally to a tumor or other pathologically affected site or organ.

FIG. 1 shows oligonucleotides useful in the present invention.

FIG. 2A shows effects of oligonucleotides (f.c. 5  $\mu$ M) on TGF- $\beta$ 2 secretion in glioma cells in 10% MEM Dulbecco medium (3 day incubation with oligonucleotides).

FIG. 2B shows effects of oligonucleotides (f.c. 5  $\mu$ M) on TGF- $\beta$ 1 secretion in PBMC in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

FIG. 3A shows effects of oligonucleotides (f.c. 5  $\mu$ M) on TGF- $\beta$ 1 secretion in PBMC in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

FIG. 3B shows effects of oligonucleotides (f.c. 5  $\mu$ M) on TGF- $\beta$ 2 secretion in glioma cells in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

FIG. 4A shows TGF- $\beta$ 1 concentration (ELISA) in glioma cells (3 day incubation with oligonucleotides).

FIG. 4B shows TGF- $\beta$ 2 concentration (ELISA) in glioma cells (3 day incubation with oligonucleotides).

FIG. 5 shows lysis of tumor-cells: LAK-Cytotoxicity, Ratio of glioma-cells/PBMC: 1:20.

FIG. 6A shows dendritic cells generated from PBMC (% of control). Cytokines: GM-CSF (400 U/ml)+IL-4 (300 U/ml).

FIG. 6B shows lysis of tumor-cells: Effects of 5  $\mu$ M VEGF-Antisense-Oligos on LAK-Cytotoxicity. Ration of tumor-cells/DC/PBMC was 1:5:20.

FIG. 7A shows effects of oligonucleotides (f.c. 5  $\mu$ M) on TGF- $\beta$ 1 secretion in PBMC in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

FIG. 7B shows effects of oligonucleotides (f.c. 5  $\mu$ M) on TGF- $\beta$ 2 secretion in tumor cells in 10% FCS RPMI 1640 medium (3 day incubation with oligonucleotides).

FIG. 8 shows lysis of tumor-cells: Effects of oligonucleotides on LAK-Cytotoxicity. Ration of tumor-cells/PBMC was 1:20.

## EXAMPLES

### Preparation of PBMC and Tumor Cells

Peripheral blood mononuclear cells (PBMC) were isolated from venous blood of healthy donors by standard Ficoll-Hypaque gradient centrifugation. Briefly, heparinized blood was mixed with equal volumes of complete medium (CM: RPMI 1640 medium supplemented with 10% (v/v) fetal calf serum and 1 mM L-Glutamine) and layered onto a Ficoll-Hypaque (Pharmacia, Uppsala, Sweden) gradient. After centrifugation at 400 g for 30 min at room temperature, PBMCs banded at the plasma-Ficoll interface were recovered, washed three times and resuspended in complete medium. Cell viability, as determined by Trypan blue exclusion, was greater than 97%.

Human glioma cell lines were established from tumor specimens of patients with anaplastic astrocytoma (WHO Grad III) or from glioblastoma (WHO Grad IV).

### Measurement of Cell Proliferation

For PBMC-proliferation assays (3H-thymidine incorporation and cell counting), freshly isolated PBMCs were cultured for 72 h in 96-well round-bottom plates (Nunc, Copenhagen, Denmark) at a final concentration (f.c.) of  $10^5$  cells/well (100  $\mu$ l CM). For cell number determination the cells were counted by hemacytometer. Cell viability was determined by trypan blue staining. Treated and untreated cells showed 95-100% viability after 72 h in vitro growth (with or without S-ODN).

For the tumor proliferation experiments  $10^4/100$   $\mu$ l glioma cells were seeded into 96-well flat-bottom plates (Nunc, Denmark) and incubated with cytokines and/or oligonucleotides. The DNA synthesis rate was measured, by a standard 3H-thymidine incorporation assay and determination of cell number was performed as described above.

### Quantification of TGF- $\beta$ 1 Protein in Culture Supernatants by Enzyme-Linked Immunosorbent Assay (ELISA)

The culture medium was harvested after 3 days, cleared of cellular components by centrifugation, filtered and stored at  $-70^\circ$  C. until processed further. TGF- $\beta$ 1 and TGF- $\beta$ 2 concentrations were measured after acidification of supernatants by TGF- $\beta$ 1 and TGF- $\beta$ 2 ELISA (R&D Systems, Minneapolis, USA) in duplicates, as recommended by the manufacturer.

FIGS. 1-4 and 7 show the effect of oligonucleotides on the TGF- $\beta$  secretion in cells. The concentration of the TGF- $\beta$  is reported as an optical density. The higher the optical density the higher is the concentration of the TGF- $\beta$ .

FIGS. 1A and 1B shows the effect of the oligonucleotides on the TGF- $\beta$  secretion. Control oligos (GAA GGA ATT ACC ACT TTC) have no effects whereas the oligonucleotides shown in the figures reduce the secretion of TGF- $\beta$ . The oligos in FIG. 1 are more effective against TGF- $\beta$ 1.

FIG. 2 shows further oligos and their effects on TGF- $\beta$  secretion. TGF- $\beta$ -14 is especially effective against the secretion of TGF- $\beta$ 1 and - $\beta$ 2.

FIG. 3 shows further oligonucleotides being effective against secretion of TGF- $\beta$ 1 and - $\beta$ 2. These oligonucleotides are more effective against TGF- $\beta$ 2 but are also effective against TGF- $\beta$ 1.

FIG. 8 shows a supra additive effect on tumor cell cytotoxicity by a combination of 2  $\mu$ M each of a TGF- $\beta$ 1 and TGF- $\beta$ 2 antisense oligonucleotide compared to a single 5  $\mu$ M dose of either oligonucleotide.

### CARE-LASS (Calcein-Release-Assay) to Measure Cytotoxic PBMC Activity

A standard calcein-release-assay (CARE-LASS assay) to determine cytotoxic activity of PBMC was employed as described by Lichtenfels, R., Biddison, W. E., Schulz, H., Vogt, A. B. and R. Martin. CARE-LASS (calcein-release assay), an improved fluorescence-based test system to measure cytotoxic lymphocyte activity. *J. Immunol. Meth.*, 172: 227-239, 1994.

### Target and Effector Cells

At the day of the assay malignant glioma were harvested, washed twice in 5% FCS/PBS and incubated with Calcein-AM (Molecular Probes, USA) for 30 min in  $37^\circ$  C. Labeled target cells were washed twice in 5% FCS/PBS, adjusted to 100 000/ml, and plated into 96-well U-shaped microtiter plates (Nunc, Denmark) at the final volumen of 100  $\mu$ l/well.

PBMC were washed with 5% FCS/PBS and adjusted to final concentration of 1-10 Mio cells/ml.

Cells were treated with cytokines and oligodeoxynucleotides as described in the individual experiments.

### Assay

To measure CTL activity effector cells were plated into 96-well U-shape microtiter plates at Target:Effector Ratios of 1:10-1:100. To measure spontaneous release and total release of calcein, wells were preloaded with 200  $\mu$ l 5% FCS/PBS or 200  $\mu$ l lysis buffer (50 mM sodium-borate, 0.1% Triton, pH 9.0) respectively. After incubating the plate for 4 h at  $37^\circ$  C. in an incubator, 100  $\mu$ l of supematans were transferred into new wells and measured employing an automated fluorescence scanner (Titertek Fluoroskan II, Germany). Both for excitation and for emission, filter settings 2 were chosen (ex 2-485

7

nm, em 2-538 nm). The percent of cytotoxicity was determined from the following equation:

$$\frac{F/CTL \text{ assay} - F \text{ spontaneous release}}{F \text{ total lysis} - F \text{ spontaneous release}} \times 100 = \% \text{ cytotoxicity}$$

In one set of experiments, glioma cells, denritic cells (DC) and PBMC were co-cultured. In these experiments DC were generated from PBMC using the cytokines GM-CSF and IL4. Cells were further treated with antisense VEGF-oligonucleotides according to the invention or with no oligonucleotides as control experiments. Tumor cells were also treated with the cytokines GM-CSF and IL4 with or without oligonucleotides.

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PBMC were only treated with oligonucleotides according to the invention, but not with the cytokines GM-CSF and IL4. oligos were used at a concentration of 5  $\mu$ M unless indicated otherwise in the descriptions in the figures.

5 The CARE-LASS (calcein-release-assay) was used to measure cytotoxic PBMC activity.

In one set of experiments glioma cells and PBMC were treated either with a single oligonucleotide or with a combination of oligonucleotides. The single oligonucleotides were given at 5  $\mu$ M concentration. In the combination experiment, each oligonucleotide was given at 2  $\mu$ M concentration. Both, PBMC and tumor cells were incubated separately with the oligonucleotide(s) for 72 h.

The CARE-LASS (calcein-release-assay) was used to measure cytotoxic PBMC activity.

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<400> SEQUENCE: 97  
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<400> SEQUENCE: 112  
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<210> SEQ ID NO 116  
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<400> SEQUENCE: 117  
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<400> SEQUENCE: 146  
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<210> SEQ ID NO 148  
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<400> SEQUENCE: 149  
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<210> SEQ ID NO 153  
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<400> SEQUENCE: 153  
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<210> SEQ ID NO 154  
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<400> SEQUENCE: 154  
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<210> SEQ ID NO 155  
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<210> SEQ ID NO 156  
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<400> SEQUENCE: 156  
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<213> ORGANISM: Homo sapiens

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<210> SEQ ID NO 161  
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<400> SEQUENCE: 161  
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<400> SEQUENCE: 162  
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<210> SEQ ID NO 163  
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<400> SEQUENCE: 165  
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<210> SEQ ID NO 166  
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<210> SEQ ID NO 167  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 167  
ccagcaatga cagc 14

<210> SEQ ID NO 168  
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<400> SEQUENCE: 168  
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<210> SEQ ID NO 169  
<211> LENGTH: 15  
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<400> SEQUENCE: 169  
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<210> SEQ ID NO 170  
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<400> SEQUENCE: 170  
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<210> SEQ ID NO 171  
<211> LENGTH: 14  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 171  
gggctggtgt ggtg 14

<210> SEQ ID NO 172  
<211> LENGTH: 16  
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<400> SEQUENCE: 172  
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<210> SEQ ID NO 173  
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<400> SEQUENCE: 173  
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<210> SEQ ID NO 174  
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<400> SEQUENCE: 174  
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<210> SEQ ID NO 175  
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<400> SEQUENCE: 175  
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<400> SEQUENCE: 176  
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<210> SEQ ID NO 177  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 177  
cccttgctc ggggg 15

<210> SEQ ID NO 178  
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<400> SEQUENCE: 178  
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<210> SEQ ID NO 179  
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<210> SEQ ID NO 181  
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<400> SEQUENCE: 181  
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<210> SEQ ID NO 182  
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<210> SEQ ID NO 183  
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<400> SEQUENCE: 183  
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<400> SEQUENCE: 184  
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<210> SEQ ID NO 185  
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<400> SEQUENCE: 185  
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<210> SEQ ID NO 186  
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<400> SEQUENCE: 186  
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<210> SEQ ID NO 187  
 <211> LENGTH: 14  
 <212> TYPE: DNA  
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<400> SEQUENCE: 187  
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<210> SEQ ID NO 188  
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<400> SEQUENCE: 188  
 gcaaagtcca gcagggc 17

<210> SEQ ID NO 189  
 <211> LENGTH: 16  
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 <213> ORGANISM: Homo sapiens

<400> SEQUENCE: 189  
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<210> SEQ ID NO 190  
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<400> SEQUENCE: 192  
gaccgtggca aagttcag 18

<210> SEQ ID NO 193  
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<400> SEQUENCE: 193  
agagaggctg accgt 15

<210> SEQ ID NO 194  
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<400> SEQUENCE: 194  
gacagagaga ggctgac 17

<210> SEQ ID NO 195  
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<400> SEQUENCE: 195  
acagagagag gctga 15

<210> SEQ ID NO 196  
<211> LENGTH: 15  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 196  
gtggacagag agagg 15

<210> SEQ ID NO 197  
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<400> SEQUENCE: 197  
caagtggaca gagagagg 18

<210> SEQ ID NO 198  
<211> LENGTH: 16  
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<400> SEQUENCE: 198  
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<210> SEQ ID NO 199  
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<400> SEQUENCE: 199  
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<210> SEQ ID NO 200  
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<400> SEQUENCE: 200  
cacctcttc ttct 14

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<400> SEQUENCE: 201  
atggatttct ttggcat 17

<210> SEQ ID NO 202  
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<400> SEQUENCE: 202  
ggatttcttt ggc 13

<210> SEQ ID NO 203  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 203  
aagttggact ctcttctc 18

<210> SEQ ID NO 204  
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<400> SEQUENCE: 204  
taagttggac tctcttct 18

<210> SEQ ID NO 205  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 205  
gacctaaagt ggactc 16

<210> SEQ ID NO 206  
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<400> SEQUENCE: 206  
tttctagacc taagttgg 18

<210> SEQ ID NO 207  
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<400> SEQUENCE: 207  
ctgatttcta gacctaag 18

<210> SEQ ID NO 208  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 208  
gaagcagtaa ttggtgt 17

<210> SEQ ID NO 209  
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<212> TYPE: DNA  
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<400> SEQUENCE: 209  
ggaatcatca tgagg 15

<210> SEQ ID NO 210  
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<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 210  
gggaatcatc atgag 15

<210> SEQ ID NO 211  
<211> LENGTH: 15  
<212> TYPE: DNA  
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 211  
ggttgtcgag ccggt 15

<210> SEQ ID NO 212  
<211> LENGTH: 17  
<212> TYPE: DNA  
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 212  
gtcctcccaa catagta 17

<210> SEQ ID NO 213  
<211> LENGTH: 14  
<212> TYPE: DNA  
<213> ORGANISM: Homo sapiens

<400> SEQUENCE: 213  
gggtcctccc aaca 14

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## 61

The invention claimed is:

1. A composition comprising

- a) an oligonucleotide having a sequence according to SEQ ID NO: 9, unmodified or having one or more modifications selected from the group consisting of phosphorothioate internucleotide linkages, methylphosphonate internucleotide linkages, phosphoramidite linkages, peptide linkages, 2'-O-modified sugar, and modified bases, which oligonucleotide reduces the expression of transforming growth factor (TGF)- $\beta_1$ , TGF- $\beta_2$ , and TGF- $\beta_3$ , in combination with
- b) at least one stimulator positively effecting an immune response by enhancing proliferation or function of immune cells and selected from the group consisting of Granulocyte Macrophage Colony Stimulating Factor (GM-CSF), Stem Cell Factor (SCF), Colony Stimulating Factor (CSF), Interferon (IFN), FMS-Related Tyrosine Kinase 3 Ligand (FLT-3-ligand), interleukin-4, interleukin-12, and interleukin 18.

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2. The composition according to claim 1, wherein the at least one stimulator is two or more stimulators.

3. The composition according to claim 1, wherein the sequence according to SEQ ID NO: 9 is unmodified.

4. The composition according to claim 1, wherein the sequence according to SEQ ID NO: 9 has one or more modifications selected from the group consisting of phosphorothioate internucleotide linkages, methylphosphonate internucleotide linkages, phosphoramidite linkages, peptide linkages, 2'-O-modified sugar, and modified bases.

5. The composition according to claim 2, wherein the sequence according to SEQ ID NO: 9 is unmodified.

6. The composition according to claim 2, wherein the sequence according to SEQ ID NO: 9 has one or more modifications selected from the group consisting of phosphorothioate internucleotide linkages, methylphosphonate internucleotide linkages, phosphoramidite linkages, peptide linkages, 2'-O-modified sugar, and modified bases.

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